

1/20

Band 55 protein

	10	20	30	40	50
ISR	MESAEELPARPALETEGLRFLHVTVGSLASYGWYVLFSCILLYIVIQK				
MOUSE	-DRD---S-@-----S-*-----I-----R				
RAT	---S-----S-----I---V-----				
HUMAN	--RQ--S-S-----T-----T---IV-----V-F--				

	60	70	80	90	100
ISR	LSVRLRALRQRQLDQADAVLEPDVAVKRQEALAAARLRMQEDLNAQVEKH				
MOUSE	--L-----ET---V-----				
RAT	--L-----E---V-----				
HUMAN	--A-----R-A-AV--V-----K--E-----				

	110	120	130	140	150
ISR	KEKLRQLEEEKRRQKIEMWDSMQEGRSYRRNPGRPQEEDGPGPSTSSSVT				
MOUSE	-----K--S-----/---I				
RAT	-----K-----/---I				
HUMAN	---K-----K--KG-AKK---S---/---L				

	160	170	180	189
ISR	RKGKSDKKPLRGNGYNPLTGEGGGTCAWRPGRRGPSGGG			
MOUSE	P-----G-----S-----			
RAT	P-----G-----			
HUMAN	/-R--R----G----S-----A S-----			

@=R or S

*=R or Q

A= mixture of A and V

Figure 1

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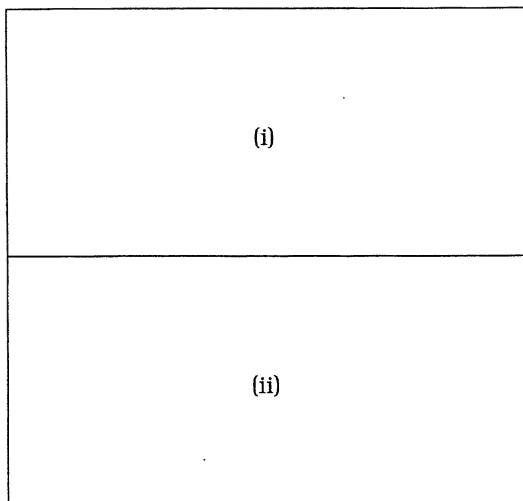


Figure 2

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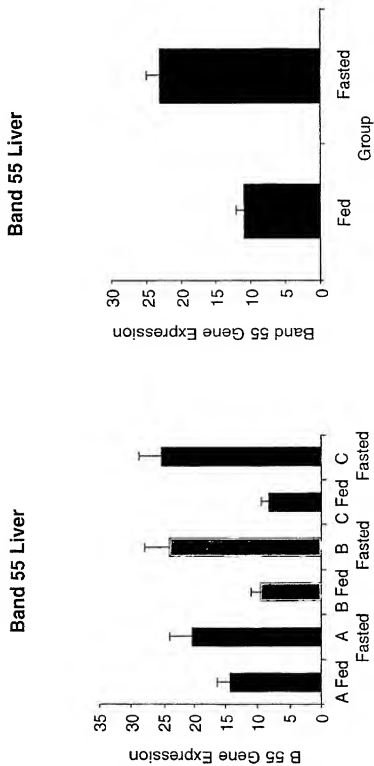
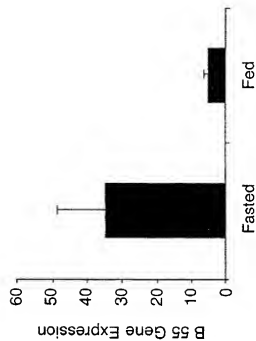


Figure 2(i)

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Band 55 Adipose Tissue



Band 55 Adipose Tissue

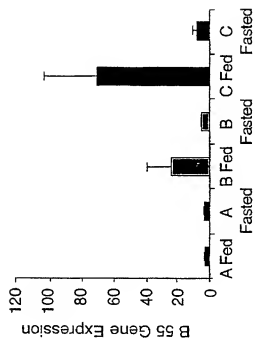


Figure 2(ii)

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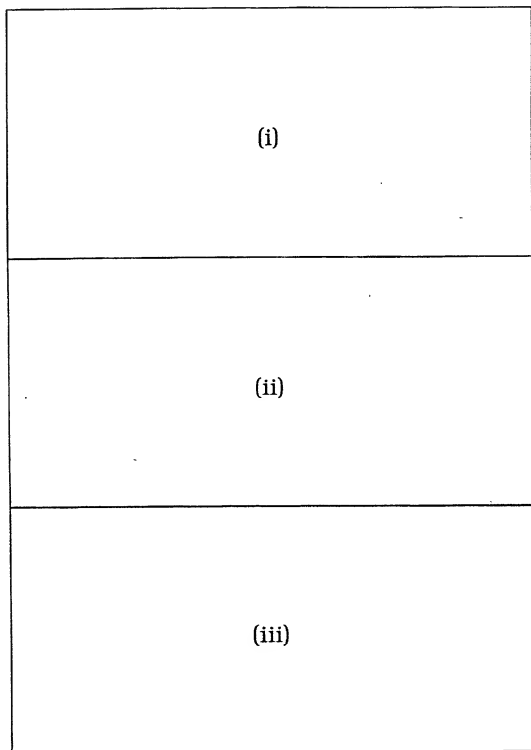


Figure 3

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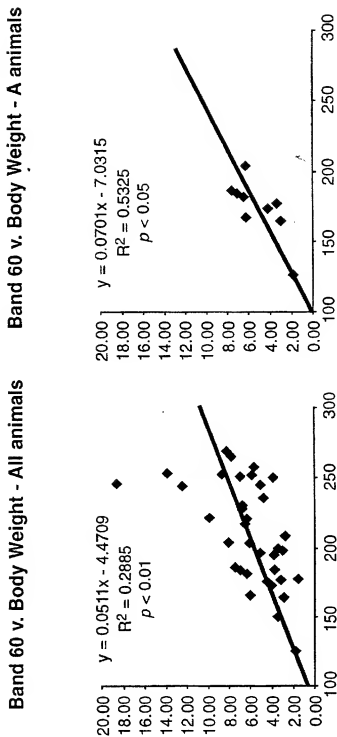


Figure 3(i)

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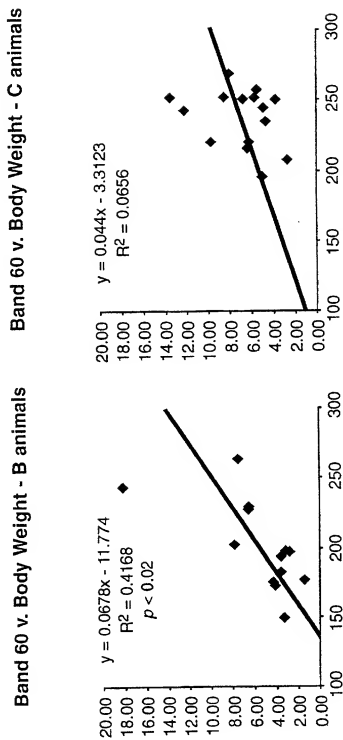


Figure 3(ii)

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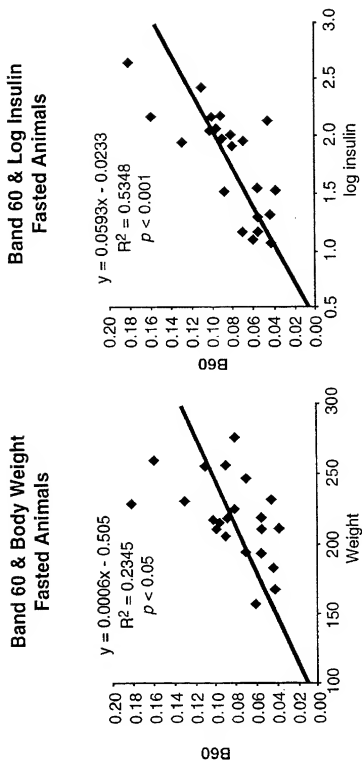


Figure 3(iii)

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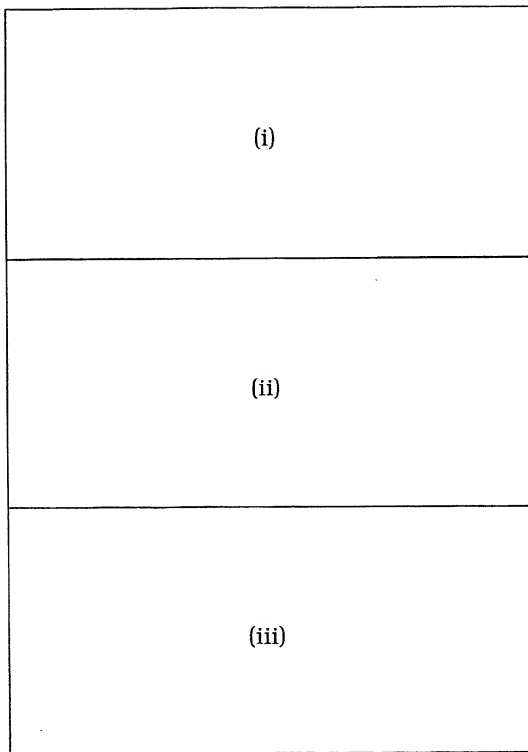


Figure 4

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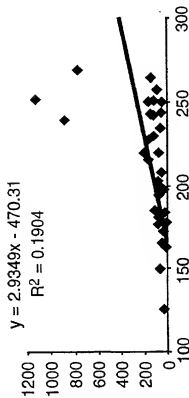
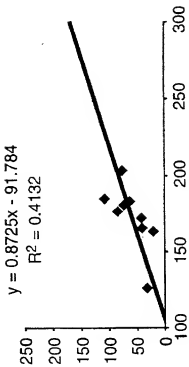
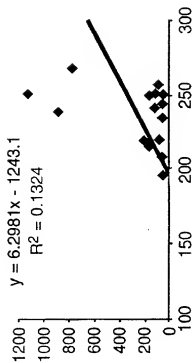
LIVER**Band 38 v. Body weight - All animals****Band 38 v. Body weight - A animals**

Figure 4(i)

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Band 38 v. Body weight - C animals



Band 38 v. Body weight - B animals

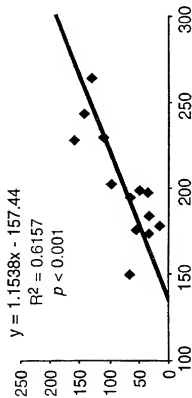
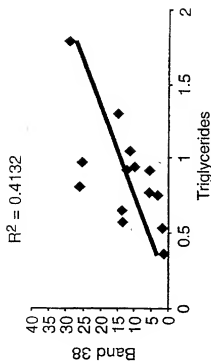


Figure 4(ii)

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Adipose tissue - Band 38 v triglycerides



Liver - Band 38 v. Triglycerides

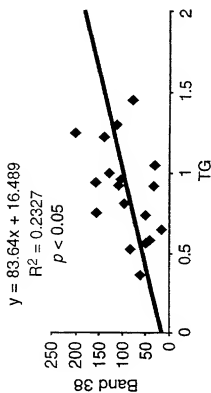


Figure 4(iii)

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Genomic structure of the human band 55 gene

Chromosome 15

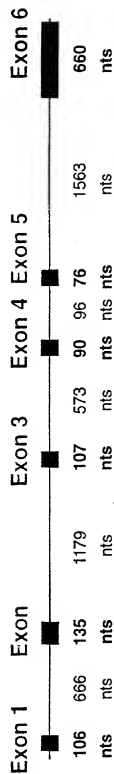


Figure 5

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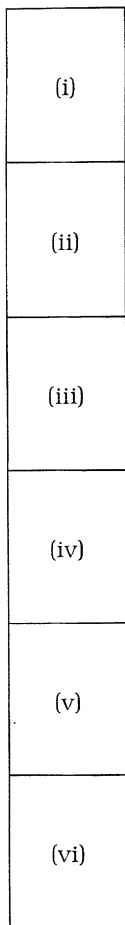


Figure 6

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1 CAGGGCTGGG CGGCGGCGGC GGC GGCGGTC ATGGAACGCC AAGAGGAGTC
↑ transcription initiation site translation start codon
start exon 1

51 TCTGTCCGCG CGGCCGCCCC TGGAGACCGA GGGGCTGCGC TTCCTGCACA

101 CCACGGTGA GTCGTTGCGG GGCAGCCGGG GCGCGCCCGC CACTTTTTCG
↑ end exon 1

151 ACGCGCAGCC ATGATGGGTG GTCGTCGCGC CGTGACACCG GGCGCCGGAG

201 CCTGGGAGGC CTGGGAACGG TCGGGCGTTG GCGCTTACGC GGACCTTGGG

251 CAGCAGGCCC GGACCTTGCG CGGAGGCTTC TCGGAGCCG CACTTCCCTG

301 GGC GGCTCGG CTGTCCCTTG TTTGCGCAAG TCTTTTTTGC GAACCAAGCC

351 CTTCTGTGG TAGTTACTGG GGTCACTCGG CCGTTGGCGT TTGCTCTGG

401 GACCCGTCAC ACACAGCCCC ATACACACTC CTGACTCCCC GCGCTGTAC

451 CCCTTTCTAT GTGGCTCTGA AAGGCCTTTG CCTTCTGAT TCAGATTAGT

501 TGCTCTTCAT TCTTCAAAC CAGTTGCTG TGCCCTCCAC ACTCTAACTG

551 CCCCCGACTC CCCAGATGGT TGGGAAGTCT CACTTCTCAG TGATCCCTGA

601 ATTGTGCGAC TTCTTGAGTT CGTGTTTTAA CGATCTACTT AGGAGGCTTT

651 TTCCTCAGCC TAGACCATGA AGGCTTTGAG GGCAGGAGTT ACACCTTTTG

701 TTTGTTGAGT CTTATGGAAA GGTCAACTAG TAGTGTCATT TTTAGTTTTT

751 TGAAAACTGT TTTTCTTTTC AGTGGGCTCC CTGCTGGCCA CCTATGGCTG
↑ start exon 2

801 GTACATCGTC TTCAGCTGCA TCCTTCTCTA CGTGGTCTTT CAGAAGCTTT

851 CCGCCCGGCT AAGAGCCTTG AGGCAGAGGC AGCTGGACCG AGCTGCGGCT

901 GCTGTGGGTT AGTGCCTGAT AACCGAAATG AAAGCGGTGG TTTTGACCT
↑ end exon 2

Figure 6(i)

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951 CCTTTATATT AAGAGTTAGT CTCTTAGTAA AAGTAAGAGG GGCCACACAG
1001 GAAGACCCTG TCTCTATTTA AAAAAAAAAA AAATAGCCGG GAGTGGCGGC
1051 ACGCACCTGT AGTCCCAGCT GCTCAGGAGG CTGAGGCGGG ATAATCACTT
1101 GAGTCCAGGG AGTCAAAGCT GCAGTGGGCT ATGCTCGGGC CACACTACAC
1151 TCCAGCCTGG GCAATTGATT GAGACCTTGT CTTTAAAAAA AAAAAAAAAA
1201 AAAAAGTAGG AAGTATATGG TTCTCGGTGG GCGCGGTGG CTCACACCTG
1251 TAATCCAGC ACTTTGGGAA GCCGAGGCAG GAGGATGACT TGAGGTACAG
1301 GGTTTCGAGAA CAGCCTGGCC AACATGGTGA AACCTGTCT CTAATAAAA
1351 TACAAATATT AGTGGGGCGT GGTGACGGGC ACCTGTAATC CCAGCTATTA
1401 GGGTGGCTGA GGCAGGAGAA ATCGCTTGAA CCTGGGAGCT GGAGATTGCA
1451 GTGAGCTGAG ATTTGCCAC TGCACTCCAG CCTGGGCAAC AGAGTGAGAC
1501 TGTCTTTTCT TTCTTTTTTT TTTTTTTTTC TATGAGATGG AGTCTAGCCT
1551 TGTTGCAAAG AGCGAGACTC TATGAGTAGA CGTTATGAAT AGAAATGAGT
1601 TCATTCTAT TCATAATGCT ATTTGGAAGG ATTTTCTTT TCTGTAGAAA
1651 CAAATACTTA AGAATCTTCT GCGCTAATTA AGGGATGGAT AATGATTAG
1701 AAAACTTTAT ATTTCTTGG TAGTCTTCCA GGATTCTAGT CAGCCTAGAG
1751 ACTGTGGGTG TCACTGAGGT ATCCAAGATG TGCTCTGTGT GGCCACTATC
1801 CCAGGCTTTA TGAATCGGAA TTGCTCAGGG GAACTCAGAA ATTGGCATT
1851 CTAACAGATT TCTGGTGATG TAGATATTC GGGCTAAAT CCGTGGCTCA
1901 GCAACAGACC CTGCCCCCT GAAGCAGTAA AATGTATGCA GAGGGGTAG
1951 GAGTACTTAT GTAAAAATAT GTTGTTTCAT TGTCTGATAT CCATACCTCT
2001 TTATACTTTT AATAATATGG ACACTCAAAA GTTTCATTTT TATATTGTAC

Figure 6(ii)

SUBSTITUTE SHEET (RULE 26)RO/AU

17/20

2051 ACAGTGCTTT ATCTCCATTT TTTTCTGACA TTTTAGAACC TGATGTTGTT
↑
start exon 3

2101 GTTAAACGAC AAGAAGCTTT AGCAGCTGCT CGACTGAAAA TGCAAGAAGA

2151 ACTAAATGCG CAAGTTGAAA AGCATAAGGA AAAACTGAAA CAAGTATGAA
↑
end exon 3

2201 CTGGTTTCAG TTTGAATGTG TGCATAGAAA TTGTCTGAGG TTTAGTGGCT

2251 AACGATGCCT GTGTCTGTGT TGTCTATAAG CTTCTAGGAC CAGGTCCTAT

2301 CCCATTAGAT TCAATAAGCA TTTCAGTTCC TACCATGTAA GTATTGGTGA

2351 TATCAAGAAG AATACACGAT TGTTAGGGAA CACTAGATGT GTGAATATAT

2401 TACCATGAAA GGTCCAGAGC ACAAAGGAG GGACAGGCTG GAGCAGGGAG

2451 CATGTGAGTG TGTGTGTGCA TGTGCCTGTG TCTTCCCCAT TACCAAAAAAT

2501 GTCCTGACAG GAGTGAGTTT CAGAAGAATG GAGTCAGTAA TCTTTTTCAT

2551 GAAACATTTT GCTTTCTTTA ATAGTGTACA AAAACCAAAG CTGCTCTATG

2601 TGAGTTAAAC TCACACTACC AGATCACAAC AGTTTTATTA ACTAAAGAAA

2651 ACGAGGGTGA AGTTTGTCTT GAAAGACATT TAAATTAAGA ATTATCAGAG

2701 TTAGCTTTGT CTTTGAGAGA AATGGCAGCT TCTGAATTCT TTCTGTAAAA

2751 TGTGATTGTT TCTCAGCTTG AAGAAGAAAA AAGGAGACAG AAGATTGAAA
↑
start exon 4

2801 TGTGGGACAG CATGCAAGAA GGAAAAAGTT ACAAAGGAAA TGCAAGAAG

2851 CCCCAGGTGA CTGGAGACCT CGGCCGGCTG GCATGCGGTA GATGAAGATT
↑
end exon 4

2901 GCCAAGTAGA ATGTTTTAAT TGCTTCTTAC ACTACTGTGT GTGTTCAAAC

Figure 6(iii)

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2951 AGGAGGAAGA CAGTCCTGGG CCTTCCACTT CATCTGTCCT GAAACGGAAA
↑
start exon 5

3001 TCGGACAGAA AGCCTTTGCG GGGAGGAGGT AAGCACCCTT GATGTCAAAT
↑
end exon 5

3051 GTTAACAGAT TTTCACACT TACAGGATAT AGTTACCTTT TAAGAACAAG
3101 ATTGTTTGTT TCTTTGTCCA TAAATTAAGA CTAATTCCTT AGGATTGTGA
3151 AGATTCAATA AAGGAAACAG ATGCAAATCA CCTCCTAGGT CCTCACTAAG
3201 TACTTAGAAG GATTGTACTT ATAGTATTCT AACTTGATCC TTCTGCAGCC
3251 CCGTAGAGGG AGAGCTAAGT AGGGTGAGGA ATTGTCTGCC AATCTTCAGA
3301 TGAGTGTCAA GGAGCTGGAA CACAGTGGTT TTGGTCTTTC TGGCTGGGAC
3351 CACCTTGTTT CTGCAAATA ACAAGGAGTA GCAGACAGAT GCTCATCCAA
3401 AGCTGCTTCC TGTGTGCAGC ACTGCCCGG GGA CTCTGGA TGATGCCACA
3451 GCAGTCTGTC TTCATCCCAT CCCTGAGAAT TTCAAATCTG GGAAGATGGG
3501 ACTCACAAC GAAAATAAGC AATCCTTGGT GATTCTGGCT AAGAGTTGCA
3551 AGTTACTGCT GAGGAAGGAA AGAACAAACA CACTAGAACA CTGTAGGAAC
3601 CAAGGCGGAA GATTTTGTAT CCTCCATAGG AGGAGAGGGG CACCGCAGAG
3651 GCCCTGATGG TGTCTTTGAG GACTGAGGAA AGACTGGGGC ATGGGCTCCA
3701 AGGCAGCAGG GCCACAGACT TGGCTGACCT TAAACGCTGA GCTGTAATCC
3751 CCTTTGTGTC AGAAGACTAA ACCTGGCTTG CTGTAGAGAA GGTGATGCAT
3801 CTGAAAGAA AATGCTATTT TTAAATGGTC CTGCCGAAG CTTATTTTTA
3851 GACACATAGA GGTGATATTT AGGAGAGGAA TGGAAATCGT AGAAGATGGA
3901 ATGCGGGTG TGCTTGCTTG CACGGCCTCT TTCAGCATCC CCAGCATTTT
3951 TGAGCTGGGA CTTTGTACTA GCCTGGCTTT ACAAATAAGG AACTGAGGC

Figure 6(iv)

19/20

4001 ACAGTGTTTA ATTGCCCAA GATTCCACTA TAAGTAAGGA GTAAAAGTAA
4051 CATTTAAGTT CTGGGTGGCC CTAGAACCCT AGCACTCAAC CAGGTTACCA
4101 GTTGTGCACT GACTTTGGGA AGCTCATGAG GGAGTGGGGT GGTGGGGGT
4151 AGGGAAGGAT ACAGAAGACC CCGTTCGAC TGGTAGAAGT GACAAGTTG
4201 ACTCTTGATT TTTTTTAATC TGTTTTCTGT AGCGTGAACA GCCCTTATTT
4251 GAATGTATGA GTTTTAGTAA GCACTGTGAT AGGAGGATTC ATATACTTAA
4301 ATCAGGCCCT CTTGAGAGAG TTTTTTGGTG ACCCTTTTGC ATGTGTTTCG
4351 GAGGTTGGGA CAAAGAAGCT GAATGACTTT TTTCCCACC AGACAATCAG
4401 TTCAAATGCG AATCACAATA TAAAGGTTTT TTTTTTTTTC ACATAGCTAA
4451 AAGGTTTTTT TAAATGTCCC TTAGGATCTG TATCTTTGCA GTGCTTTGCG
4501 TGTCACTCTC ATAATTTTAT TGTGGATATA CAATGTTCCC AGATTTTCAG
4551 ATTTTATCA ATACTGTTGT GCTGCTTTTC TGTCTCCCA GGTATAACC
↑
start exon 6
4601 CGTTGTCTGG TGAAGGAGGC GGAGCTTGCT CCTGGAGACC TGGACGCAGA
4651 GGCCCGTCAT CTGGCGGATG AGGCTAAGAA TCTTGTTAGT GTCACTTTTG
↑
translation stop codon
4701 ACATTAGCAA GATGAACCCT TAACCCTCGA TTCAATTGCC TTACGCACGC
4751 TTTTCACAGT GACTAGCCAA GGGGAGGTGG GGTGATTTC TGTCCTAAC
4801 TACACCTGCA TATGTCAGGG CTCCAGTCAG CAAAAGGTAT AGATGTTGCC
4851 TCTAGGCATG AGGTCATTGG TCACATTCTA CTTGGAGACA GTGATTGCAT
4901 TCATTGATTT CATGGTTAAT TGCTAGTTGG TAGGTAAAGG CCTCTAGATG
4951 ATTAGCAATC TTGATAAAAG AGGCCTAGTA ATGTTCTTTT GAGGTTAGAA
5001 ATCCTTGCTG CTAGGACAGT CTCTGTGACA GGTGCGTTG AATGATGTCT

Figure 6(v)

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5051 TCCTTATCAA TGGTGAGCCC ACCAGTGAGG ATTACTGATG TGGACAGTTG
5101 ATGGGGTTTG TTTCTGTATA TTTATTTTTA TGTACAGAAC TTTGTAAAAA
5151 CGAAACTATT TAAAAACAA GAATAACATT TTTAGCATCT TTATTCAAGG
5201 AGATTTATGG ACTTCAATTT GTCTATCAAA CATTAAATAG CTTTTTATTA
5251 C
↑
transcription termination site
end exon 6

Figure 6(vi)